

INFORMATION DISCLOSURE STATEMENT	Atty. Docket No.: 265.0041 0101	Serial No.: 10/580,979
	Applicant(s): Lemon et al.	Confirmation No.: 9290
	371(c) Filing Date: April 9, 2007	Group: 1648
	Information Disclosure Statement filed:	December 20, 2009

U.S. PATENT DOCUMENTS

Examiner Initial	Copy Enclosed	Document Number	Date	Name	Class	Subclass	Filing Date If Appropriate
		<i>None</i>					

U.S. PATENT APPLICATIONS BY SERIAL NUMBER

Examiner Initial	Copy Enclosed	Document Number	Filing Date	Name	Class	Subclass
		<i>None</i>				

FOREIGN PATENT DOCUMENTS

Examiner Initial	Copy Enclosed	Document Number	Date	Country	Class	Subclass	Translation	
							Yes	No
	X	WO 2004/055216 A2, A3	07/01/04	WIPO				

OTHER DOCUMENTS (Including Authors, Title, Date, Pertinent Papers, etc.)

Examiner Initial	Copy Enclosed	Document Description
	X	Adams et al., "Complete Coding Sequence of Hepatitis C Virus Genotype 6a," <i>Biochemical and Biophysical Research Communications</i> , 1997; 234:393-396.
	X	Bressanelli et al., "Crystal structure of the RNA-dependent RNA polymerase of hepatitis C virus," <i>PNAS</i> , 1999 November 9; 96(23):13034-13039.
	X	Bressanelli et al., "Structural Analysis of the Hepatitis C Virus RNA Polymerase in Complex with Ribonucleotides," <i>Journal of Virology</i> , 2002 April; 76(7):3482-3492.
	X	Chamberlain et al., "Complete nucleotide sequence of a type 4 hepatitis C virus variant, the predominant genotype in the Middle East," <i>Journal of General Virology</i> , 1997; 78:1341-1347.

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*Examiner: Initial if citation considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.	

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Examiner Initial	Copy Enclosed	Document Description
	X	Cheney et al., "Mutations in NS5B Polymerase of Hepatitis C Virus: Impacts on <i>in Vitro</i> Enzymatic Activity and Viral RNA Replication in the Subgenomic Replicon Cell Culture," <i>Virology</i> , 2002; 297:298-306.
	X	Love et al., "The Crystal Structure of Hepatitis C Virus NS3 Proteinase Reveals a Trypsin-like Fold and a Structural Zinc Binding Site," <i>Cell</i> , 1996 October 18; 87:331-342.
	X	Love et al., "Crystallographic Identification of a Noncompetitive Inhibitor Binding Site on the Hepatitis C Virus NS5B RNA Polymerase Enzyme," <i>Journal of Virology</i> , 2003 July; 77(13):7575-7581.
	X	Marzio et al., "HIV-1 Tat transactivator recruits p300 and CREB-binding protein histone acetyltransferases to the viral promoter," <i>Proc. Natl. Acad. Sci. USA</i> , 1998 November; 95:13519-13524.
	X	Simmonds et al., "A Proposed System for the Nomenclature of Hepatitis C Viral Genotypes," <i>Hepatology</i> , 1994; 19:1321-1324.
	X	Simmonds et al., "Evolutionary analysis of variants of hepatitis C virus found in South-East Asia: comparison with classifications based upon sequence similarity," <i>Journal of General Virology</i> , 1996; 77:3013-3024.
	X	Simmonds et al., "Consensus Proposals for a Unified System of Nomenclature of Hepatitis C Virus Genotypes," <i>Hepatology</i> , 2005; 42(4):962-973.
	X	Smith et al., "Characteristics of Nucleotide Substitution in the Hepatitis C Virus Genome: Constraints on Sequence Change in Coding Regions at Both Ends of the Genome," <i>J. Mol. Evol.</i> , 1997; 45:238-246.
	X	Yan et al., "Complex of NS3 protease and NS4A peptide of BK strain hepatitis C virus: A 2.2 Å resolution structure in a hexagonal crystal form," <i>Protein Science</i> , 1998; 7:837-847.

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